

☐ 1: GI "3650064" [GenBank] Homo sapiens chromosome 17,... Related Sequences.

LOCUS AC004099 136222 bp FFI06-JUN-2000 DNA DEFINITION Homo sapiens chromosome 17, clone HCIT421KZ4, complet⇒ sequence. AC004099 ACCESSION AC004099.1 GI:3650064 VERSION KEYWORDS HTG: HOURCE human. ORGANIBM History Batcheria Eukaryota; Metažoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. REFERENCE 1 (bases 1 to 136222) Birren, B., Linton, L., Nusbaum, C. and Lander, E. AUTHORS TITLE Hemo sapiens chromosome 17, clone HCIT421K24 JOURNAL. Unpublished REFERENCE 2 (bases 1 to 136:22) AUTHORS Birren, B., Fasman, E., McKernan, K., Nusbaum, C., Richardson, P., Lander, E., Allen, N., Baker, J., Baldwin, J., Barna, N., Beckerly, R., Boutwell, C., Byrne, S., Cantu, C., Castle, A., Cooke, F., Daly, M.J., Depayre, E., Devon, E., Dewar, K., DuRette, B., Etemadi, S., Ferreira, P., Forrest, C., Gage, D., Gardyna, S., Gensheimer, S., Geraigery, K., Gilmartin, T., Gray, D., Hagos, B., Harris, K., Horton, L., Howland, J.C., Hui, L., Jacotot, L., Linton, L., MacKenzie, J., Marquis, N., McEwan, P., McGurk, A., Meldrim, J., Molla, M., Morris, W., Morrow, J., Nachman, A., Naylor, J., O'Connor, T., Favlin, B., Peterson, K., Ranganath, S., Riley, R., Roberts, D., Rollins, G., Rossello, R., Roy, A., Shyam, R., Scohoo, S., Stilwell, J., Stone, C., Strickland, C., Sydney, E., Tang, L., Vassiliev, H., Vo, A., Widner, A., Wheeler, J., Wu, Y., Ye, W.J., Zemtseva, I., Zhao, J. and Booty, M. Lirect Submission TITLE RATEMAL Palmitted (01-FER-1998) Whiteher: Institute MIT Center to selecte Ferranch, 3.0 Charles Street, Cambridge, MA (2141, USA FFFEEFICE 4. (ballets 1 to 136222) AUTHORG Birren, B., Fasman, F., Linton, L., Nusbaum, C., Lander, E., Allen, N., Baker, J., Baldwin, J., Barna, N., Beckerly, R., Benn, J., Poatin, C., Boutwell, C., Brown, A., Byrne, S., Cantu, C., Castle, A., Cerny, J.,

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Cooke, P., Popayrë, E., Devon, K., Dewar, K., Donelan, L., DuRette, R.,

Erremadi, N., Ferreira, P., FitzHuch, W., Ferrest, C., Funke, F., Sere, St., Serbara, N., Serbara, R., Gilmartin, T.,

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Wagner, A., Wang, B., Wheeler, J., Wu, Y., Ye, W.J., Zhao, J. and Zody, M.
  TITLE
             Direct Submission
             Submitted (09-MAY-1998) Whitehead Institute/MIT Center for Genome
   JOURNAL.
             Research, 320 Charles Street, Cambridge, MA 02141, USA
                (bases 1 to 1/622..)
REFERENCE
  AUTHORS
             Firren, B., Linton, L., Nusbaum, C., Lander, E., Allen, N., Anderson, M.,
             Baker, J., Baldwin, J., Barna, N., Beckerly, R., Benn, J., Boutwell, C.,
             Erown, A., Castle, A., Cerny, J., Colangelo, M., Collins, S.,
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             Faratas, A., Lehoczky, J., Macdonald, P., Marquis, N., McEwan, P.,
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             Vassiliev, H., Vo, A., Wagner, A., Wheeler, J., Wu, Y., Wyman, D.,
             Ye, W.J., ?hac, J. and Sody, M.
  TITLE
            Direct Sulmission
  JOURNAL
            Submitted (25-SEF-1993) Whitehead Institute/MIT Center for Geneme
            Fesearch, 320 Charles Street, Cambridge, MA 02141, USA
PEFERENCE
            5 (bases 1 to 136212)
  AUTHORS
            Firren, B., Linton, L., Musbaum, C., Lander, E., Allen, N., Anderson, M.,
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            Frown, A., Castle, A., Jerny, J., Colangelo, M., Collins, S.,
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            Ye, W.J., Chao, J. and Cody, M.
  TITLE
            Direct Submission
  JOURNAL
            Submitted (06-JUN-2000) Whitehead Institute/MIT Center for Genome
            Fesearch, 320 Charles Street, Cambridge, MA 02141, USA
COMMENT
            On Sep 25, 1998 this sequence version replaced gi:3126780.
            All repeats were identified using RepeatMasker: Smit, A.F.A. &
            Green, P. (1996-1997)
            http://ftr.genome.washington.edu/FM/RepeatMasker.html.
                      Location Qualifiers
FEATURES
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     repeat region
                     637..945
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                     complement (1015..1189)
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                   'rrt family "AluCy"
                   -- mplement (14-5..1+./4)
"ipt family "Aludg"
                   - mplement (1626..1667)
/rpt family "(TAA)n"
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                    ipt family "AluSx"
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                    drpt ramily="L1MC4"
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                    'ipt family "AluSg"
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                    'rpt family:"AluSp"
                    complement (3503..3614)
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                    /rpt_family="AluSc"
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                   -complement (3692..4000)
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                   complement (4002..4323)
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                   complement (4479..4663)
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                   complement (5338..5392)
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                   complement (7291..7526) .
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                 /rpt family="L1MC5"
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                  cpt_family "LIMED"
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                 /note="Probably 28 As. Three clones read 28 As, one clone
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                 complement (12093...12213)
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                  'rp'_family-"L1MC5"
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                  'rp' family="AT rich"
                 complement (12230..12396)
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                  rpt_family="AluJb"
                 complement (12414..12520)
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                  :p' famely="LIMD'"
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                 .rpt family="AluJo"
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                 complement (13112...13455)
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                  rpt_family="LIMD5"
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                 /rpt family="(CAAA)n"
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                  rpt family="AluSx"
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                 complement (38732...38932)
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                  rpt family "Alua";"
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                  "rpt family "(TAAA)n"
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                 complement (42118..42275)
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                 /rpt_family:"FRAM"
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t jed todila.
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                 46949..47244
                 /rpt_family "Aludo"
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                 complement (47385...47555)
                 /rpt family:"MIE"
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                48501..48796
                 /rpt family "AbuSp"
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                 60134..60415
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                 51505..61813
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repeat region
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                 64291..64409
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                 64410..64436
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                 Jipt family "AluJb"
                 complement (64137..64803)
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                 rpt_family "AluJb"
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                 rpt_family:"AluSp"
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                ript family "AluY"
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                   complement (74611..74638)
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                  rpt family "AluSx"
199 4 1911 1
                  complement (83761..84031)
                  /rpt family "AluJb"
                  complement (84036..84257)
                  orpt family "Aloce x"
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/rpt family="AluSx"
  1 - 3' 1 - 1 - 1
                 complement (36634..86884)
                 .rpt family "AluSx"
 complement (37218...87329)
                 .rpt_family="LTR15A"
the at the first Mr.
                 #7409..87572
                 .rpt_family="MER5A"
                 67603..97835
gerrar aregina.
                 .rpt_family="AluSx"
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                 87886..37936
                 rpt_family="(TA)n"
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                 57958..38243
                 .rp*_family="LIME2"
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                 88256..88281
                 apt family="AT rich"
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of small region.
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                  ipt family="(TA)n"
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                - omplement (88445..88472)
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                rpt family "AluSx"
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                complement(90738..91067)
                 rpt_family="AluJb"
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                 > rpt family="POLY A"
                 complement (94612...94907)
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                complement (95810..96030)*
                 crt family="AluJb"
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repeat region
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                complement (100171...100477)
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                complement (101611..101782)
                /rpt family="AluJb"
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                complement (101786..102086)
                /rpt_family="AluSg"
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 Tarfair Inch. 1
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 tessor and a
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 de timati de qui ti
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                                        complement (107344...107524)
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                      /rpt_family "L2"
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      reference region.
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                      complement (1234°2...123761)
                      ript family="Alusq"
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      1 1 1 1 1
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                      127683..128002
     " National Legist.
                      First family="AloSq"
      return real d
                      128003..128297
                      'rpt family "AluSp"
                     complement (1:8743..129036)
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                       rpt family="AluJb"
     133958..1.3978
                      .rpt_tamily <"AT_rich"
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     ing raid indica
                      .rpt family "AT rich"
     tepeat region
                     complement (134127...134173)
                      rpt family="MEF4B"
     repeat region
                     complement (134278..134578)
                      /rpt family="AluSq"
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                      rpt family "AluSq"
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4021 ficknegačný hyfrettycte flatencecky jetygdaytyc agtygogtyn tetepytten
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4441 tyadaaygga agaaacaatt tittgtgggtg ataatggtaa taataatact aattataagt
4501 acttgetety tuccaggeae cactttyayt actttacaty aattaaccet taaatcatea
4561 ctqtaatcct atgaaqtacc tatgattagt atcoccattt tatagatgaa qaaatitagg
46-21 othagagagg thaagthact fightbagggh bababaggan ntabbatath thatbagatb
4631 taagacarea tigattataa gaagcaeete aatticagaa aligitaaaat giaaaaetgg
4741 ctgatgaaag ttgtgatgta ctttcgatta taagatgaat cttgatttca gagagttttt
4801 ttattttttt gagadagggt ttdadtgtgt detdtaggdf ggagfgdagf ggd+tgatda
4861 tggctcaatg ceacetetat ettecagget taagtgatee ttetacetea getteceaag
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5011 toctateacc trajectore againstag gattactors ggcotactte agggaetttt
5101 taftittett taalaaaaaa falagaatii itiittagaga tygggtetty eteigetytt
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522/1 gatoctorto octrăgecty crotgagtaa ctaggictat gggcacaago cacaatgocc
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5401 agttgtofta gaalaaagta fittitticag acttgtiitti totovatgag gyocantgaa
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5631 agaaaggtga atcacaagat caggagtteg aggecagect ggccaaggtg gtgaaateee
5641 gtototgata aaaatacgaa aattagotga gtgttgtgoo ggotgootgt aatoocagot
5701 actogaqagg ofgaagcagg agaatcaact tgaacotggj aggosgsgyt tgcsytgagc
5761 tgagatggeg coactotage etgggtgaaa gagcaagaet eegteteaga aaaaaaaaa
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7501 gaaftcactg cottaaggit traggaagga tgocactgit tigalcaggg catgitetec
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   8581 atttratita ittatigaaa aaaaaaatag agacagggit tetechigii geocaggdag
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71.001 aacaagtaat gtgtttaagt acaactyttg gtaaagtttt aataatootg atagaoggtg
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91031 geotecraaa ytgotgggat tärätgtaty agecaeratg negagecare eccencerae
91081 cocactitti ittitttttt tittägakaat enceagagag otgaattita kattgattto
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47561 gtoreggga gatqqqcaqi itottotoca tiaaanqcaq macatomytt tamtoqcait
 97621 throspittt fatt milite igabattane alabitaaatu pagataaatg taftriabit
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 97861 agagataagt gagelagett etggtgebåd gtgbagtttg ggtgggabaa tittabetga
 979.11 agtöctgcag actggcogtg gggatgacgg gggcoccyth cettgeggtg tinticatat
 97931 egatteaggg Hagegaactg tgeagtgete geattetatt tattggteee tgteaceeeg
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 98161 gegtgataet ggtgeectee alaggeeage filjaaastgt etgtttotga gtigaalgee
 932.11 tgccagaaat aaacagagat qfggggcaag tcaggctttt tctgcatctc agagcttttc
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 98581 ageographic tractacage atgeaatest ttacastest gragerace acaagestat
 98641 catqacaaaa aattacaaqa qaqttaqaqa auqaqaqaqt tagaaaacga qaqaqttaga
 98701 qaaqqaqaqa taaqaastet tagqqqtetaa tqysattqac ostaatqeet aatqaateta
 93761 cartatoana aagtoobaac aggtatggay tiaaanaaaa atgcotogaa atgaatgygo
 988. I attabacada taaqaattat thodagatga adaqqottat dagtaagota dagtoagoda
 OwwhI tgqaaaagta gagtocaaac etcaaccant testggggtt agttttstaa tatgsactys
 9m441 maataaacaa aactotaaaa oocaatttgo tgottgtotg ttaaaaatggo taaarggota
 9 KHOL datodrąmaą otottijato todatotyty tytbačtaca mottopagyo tiedagamyt
 9.40 \pm 1 tycticattot igticaccocat ticaticadaja galaaaataad tycaacacca galagecaatt
 9 HT1 atgtganada agdatgdana taagadagaa qtggadddad adagadadat taggagadto
 9.4%1 catchaacht tatygghach aaactyacto agtgaaggat toatgattigg aacagtteag
 99241 gatggcaatg gatagastca tobaatttgb betebabast baaaattaaa atggagabgb
 9.301 taagataagg tyfnactygt tratactiit itaggatgit cilitigeagg cilicictict
 9939-1 tootgaaata acgittgaaac cacaqqicaa caacattooc agaacagggo otottgooto
 994. I canamarate thincaggit caccettite ecotgacete eccaetatge araggytyet
 HMF1 geaggratga gageegegte cattecetes acagetross igetgonete togagggaat
 99541 gtqtetqete etaleateng ettittaggne itgaljaaast beebacetet gytenenaa.
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 Prof. 1 tetgosaggs ggetgegtga ettacooggs tootsgagte casatteaag aggeacaete
 99\%.1 bacaggerang objectings a tittinative cargoogtia catacaging jaanaateer
997M1 gegagtitte atetggetgt gaggaegaba agabaaatgg gtaagaajte babaagteba
97841 cacquights actteacted stoatcacts accessagea ggostagogg caagtacagt
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100% 61 qqqaqqqaaq qatattotat qoaqaacaaq caqeacagge aaaggetetg agetaaagat
100621 a
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